CRF Problem Report

The Scientific and Technical Information Center (STIC) experienced a problem when processing the following computer-readable form (CRF):

Application Serial Number: _09/743347_

Filing Date: _1/08/01_

Date Processed by STIC: _09/24/01_

STIC Contact: Mark Spencer, 703-308-4212

Nature of Problem:

The CRF (was):

____ (circle one) Damaged or Unreadable (for Unreadable, see attached)

____ Blank (no files on CRF) (see attached)

____ Empty file (filename present, but no bytes in file) (see attached)

____ Virus-infected. Virus name: _____ The STIC will not process the CRF.

____ Not saved in ASCII text

____ Sequence Listing was embedded in the file. According to Sequence Rules, submitted file should **only** be the Sequence Listing.

✓ Did not contain a Sequence Listing. (see attached sample)

____ Other:

PLEASE USE THE CHECKER VERSION 3.0 PROGRAM TO REDUCE ERRORS. SEE BELOW FOR DETAILS:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

http://www.uspto.gov/web/offices/pac/checker

DI ...-ñ   > _

———————————————————
———————————————————
———————————————————
———————————————————
———————————————————
———y—__————

———————————————————
———————————————————
———————————————————
———————————————————
———————————————————————————Root
Entry   `+"¡€I □
>DEST
rrf A_‾   PerfectOffice_MAIN
 & ‾ ‾   5 Perfect
Office_OBJECTS   ,
   ...$f A ...$f A
———————————————————————————

File Unreadable

Encrypted text
file unintelligeable

Actual file Contents as of

10/16/01   1:50 pm

MH
Morram Harper

DI ...-ñ   >   _~

#7

SEQUENCE LISTING

<110> Korneluk, Robert G.
      Holcik, Martin
      Liston, Peter

<120> XIAP IRES AND USES THEREOF

<130> 07891/021003

<140> 09/743,347
<141> 2001-01-08

<150> PCT/IB99/01415
<151> 1999-07-22

<150> 09/121,979
<151> 1998-07-24

<150> 09/332,319
<151> 1999-06-14

<160> 30

<170> FastSEQ for Windows Version 4.0

<210> 1
<211> 295
<212> DNA
<213> Mus musculus

<400> 1
atgtgtttgg cattatgtga agcccaaaca ctaaaaaagg agaacaaaca aaagcgcaga 60
ctttaaaact caagtggttt ggtaatgtac gactctactg tttagaatta aaatgtgtct 120
tagttattgt gccattattt ttatgtcatc actggataat atattagtgc ttagtatcag 180
aaatagtcct tatgctttgt gttttgaagt tcctaatgca atgttctctt tctagaaaag 240
gtggacaagt cctattttcc agagaagatg acttttaaca gttttgaagg aacta   295

<210> 2
<211> 299
<212> DNA
<213> Homo sapiens

<400> 2
ttttattctg cctgcttaaa tattactttc ctcaaaaaga gaaaacaaaa atgctagatt 60
ttactttatg acttgaatga tgtggtaatg tcgaactcta gtatttagaa ttagaatgtt 120
tcttagcggt cgtgtagtta tttttatgtc ataagtggat aatttgttag ctcctataac 180
aaaagtctgt tgcttgtgtt tcacattttg gatttcctaa tataatgttc tcttttttaga 240
aaaggtggac aagtcctatt ttcaagagaa gatgactttt aacagttttg aaggatcta 299

<210> 3
<211> 711
<212> DNA
<213> Homo sapiens

<400> 3

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atgacgggtt atgaagcccg gctcattact tttgggacat ggatgtactc cgtcaacaaa 60
gagcagcttg caagagctgg attttatgct ataggtcaag aggataaagt acagtgcttt 120
cactgtggag gagggctagc caactggaag cccaaggaag atccttggga acagcatgct 180
aaatggtatc caggttgcaa atatctgcta gaagagaagg gacatgaata tataaacaac 240
attcatttaa cccgttcact tgagggagct ctggtacaaa ctaccaagaa aacaccatca 300
ctaactaaaa gaatcagtga taccatcttc cctaatccta tgctacaaga agctatacga 360
atgggatttg atttcaagga cgttaagaaa ataatggagg aaagaattca aacatctggg 420
agcaactata aaacgcttga ggttcttgtt gcagatctag tgagcgctca gaaagacact 480
acagaaaatg aattgaatca gacttcattg cagagagaaa tcagccctga gagccgcta 540
aggcgtctgc aagaggagaa gctttgtaaa atctgcatgg acagatatat cgctgttgtt 600
tttattcctt gtggacatct ggtcacttgt aaacaatgtg ctgaagcagt tgacagatgt 660
cccatgtgca gcgcggttat tgatttcaag caaagagttt ttatgtctta a        711
```

<210> 4
<211> 236
<212> PRT
<213> Homo sapiens

<400> 4
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Met Thr Gly Tyr Glu Ala Arg Leu Ile Thr Phe Gly Thr Trp Met Tyr
 1               5                  10                  15
Ser Val Asn Lys Glu Gln Leu Ala Arg Ala Gly Phe Tyr Ala Ile Gly
             20                  25                  30
Gln Glu Asp Lys Val Gln Cys Phe His Cys Gly Gly Gly Leu Ala Asn
         35                  40                  45
Trp Lys Pro Lys Glu Asp Pro Trp Glu Gln His Ala Lys Trp Tyr Pro
     50                  55                  60
Gly Cys Lys Tyr Leu Leu Glu Glu Lys Gly His Glu Tyr Ile Asn Asn
65                  70                  75                  80
Ile His Leu Thr Arg Ser Leu Glu Gly Ala Leu Val Gln Thr Thr Lys
                 85                  90                  95
Lys Thr Pro Ser Leu Thr Lys Arg Ile Ser Asp Thr Ile Phe Pro Asn
            100                 105                 110
Pro Met Leu Gln Glu Ala Ile Arg Met Gly Phe Asp Phe Lys Asp Val
            115                 120                 125
Lys Lys Ile Met Glu Glu Arg Ile Gln Thr Ser Gly Ser Asn Tyr Lys
        130                 135                 140
Thr Leu Glu Val Leu Val Ala Asp Leu Val Ser Ala Gln Lys Asp Thr
145                 150                 155                 160
Thr Glu Asn Glu Leu Asn Gln Thr Ser Leu Gln Arg Glu Ile Ser Pro
                165                 170                 175
Glu Glu Pro Leu Arg Arg Leu Gln Glu Glu Lys Leu Cys Lys Ile Cys
            180                 185                 190
Met Asp Arg Tyr Ile Ala Val Val Phe Ile Pro Cys Gly His Leu Val
        195                 200                 205
Thr Cys Lys Gln Cys Ala Glu Ala Val Asp Arg Cys Pro Met Cys Ser
    210                 215                 220
Ala Val Ile Asp Phe Lys Gln Arg Val Phe Met Ser
225                 230                 235
```

<210> 5
<211> 12
<212> DNA
<213> Homo sapiens

<400> 5
```
tgttctctttt tt                                                  12
```

-2-

```
<210> 6
<211> 12
<212> DNA
<213> Homo sapiens

<400> 6
aaaaagagaa ca                                                      12

<210> 7
<211> 15
<212> DNA
<213> Homo sapiens

<400> 7
gtttcttagc ggtcg                                                   15

<210> 8
<211> 15
<212> DNA
<213> Homo sapiens

<400> 8
cgaccgctaa gaaac                                                   15

<210> 9
<211> 15
<212> RNA
<213> Homo sapiens

<400> 9
cgaccgcuaa gaaac                                                   15

<210> 10
<211> 12
<212> RNA
<213> Homo sapiens

<220>
<221> variation
<222> (1)...(1)
<223> Wild-type polypyrimidine tract.

<400> 10
uguucucuuu uu                                                      12

<210> 11
<211> 12
<212> RNA
<213> Homo sapiens

<220>
<221> variation
<222> (1)...(12)
<223> Positions 1 and 3-12 are mutated.

<400> 11
agaagagaaa aa                                                      12
```

```
<210> 12
<211> 12
<212> RNA
<213> Homo sapiens

<220>
<221> variation
<222> (1)...(12)
<223> Positions 1-2, 7, and 8-12 are mutated.

<400> 12
cuuucuuucc cc                                                    12

<210> 13
<211> 12
<212> RNA
<213> Homo sapiens

<220>
<221> variation
<222> (1)...(2)
<223> Positions 1-2 are mutated.

<400> 13
aauucucuuu uu                                                    12

<210> 14
<211> 12
<212> RNA
<213> Homo sapiens

<220>
<221> variation
<222> (3)...(4)
<223> Positions 3-4 are mutated.

<400> 14
ugaacucuuu uu                                                    12

<210> 15
<211> 12
<212> RNA
<213> Homo sapiens

<220>
<221> variation
<222> (5)...(6)
<223> Positions 5-6 are mutated.

<400> 15
uguuaacuuu uu                                                    12

<210> 16
<211> 12
<212> RNA
<213> Homo sapiens

<220>
```

<221> variation
<222> (7)...(8)
<223> Positions 7-8 are mutated.

<400> 16
uguucuaauu uu   12

<210> 17
<211> 12
<212> RNA
<213> Homo sapiens

<220>
<221> variation
<222> (9)...(10)
<223> Positions 9-10 are mutated.

<400> 17
uguucucuaa uu   12

<210> 18
<211> 12
<212> RNA
<213> Homo sapiens

<220>
<221> variation
<222> (11)...(12)
<223> Positions 11-12 are mutated.

<400> 18
uguucucuuu aa   12

<210> 19
<211> 268
<212> DNA
<213> Homo sapiens

<400> 19
tattctgcct gcttaaatat tactttcctc aaaaagagaa aacaaaaatg ctagatttta 60
ctttatgact tgaatgatgt ggtaatgtcg aactctagta tttagaatta gaatgtttct 120
tagcggtcgt gtagttattt ttatgtcata agtggataat ttgttagctc ctataacaaa 180
agtctgttgc ttgtgtttca cattttggat ttcctaatat aatgttctct ttttagaaaa 240
ggtggacaag tcctattttc aagagaag   268

<210> 20
<211> 267
<212> DNA
<213> Mus musculus

<400> 20
atgtgtttgg cattatgtga agcccaaaca ctaaaaaagg agaacaaaca aaagcgcaga 60
ctttaaaact caagtggttt ggtaatgtac gactctactg tttagaatta aaatgtgtct 120
tagttattgt gccattattt ttatgtcatc actggataat atattagtgc ttagtatcag 180
aaatagtcct tatgctttgt gttttgaagt tcctaatgca atgttctctt tctagaaaag 240
gtggacaagt cctattttcc agagaag .   267

<210> 21

-5-

```
<211> 163
<212> DNA
<213> Homo sapiens

<400> 21
aattagaatg tttcttagcg gtcgtgtagt tattttttatg tcataagtgg ataatttgtt 60
agctcctata acaaaagtct gttgcttgtg tttcacattt tggatttcct aatataatgt 120
tctcttttta gaaaaggtgg acaagtccta ttttcaagag aag        163

<210> 22
<211> 162
<212> DNA
<213> Mus musculus

<400> 22
aattaaaatg tgtcttagtt attgtgccat tattttttatg tcatcactgg ataatatatt 60
agtgcttagt atcagaaata gtccttatgc tttgtgtttt gaagttccta atgcaatgtt 120
ctctttctag aaaaggtgga caagtcctat tttccagaga ag        162

<210> 23
<211> 103
<212> DNA
<213> Homo sapiens

<400> 23
agctcctata acaaaagtct gttgcttgtg tttcacattt tggatttcct aatataatgt 60
tctcttttta gaaaaggtgg acaagtccta ttttcaagag aag        103

<210> 24
<211> 102
<212> DNA
<213> Mus musculus

<400> 24
agtgcttagt atcagaaata gtccttatgc tttgtgtttt gaagttccta atgcaatgtt 60
ctctttctag aaaaggtgga caagtcctat tttccagaga ag        102

<210> 25
<211> 83
<212> DNA
<213> Homo sapiens

<400> 25
gttgcttgtg tttcacattt tggatttcct aatataatgt tctcttttta gaaaaggtgg 60
acaagtccta ttttcaagag aag        83

<210> 26
<211> 83
<212> DNA
<213> Mus musculus

<400> 26
agtccttatg ctttgtgttt tgaagttcct aatgcaatgt tctctttcta gaaaaggtgg 60
acaagtccta ttttccagag aag        83

<210> 27
<211> 129
<212> DNA
```

    <213> Homo sapiens

    <400> 27
    aattagaatg tttcttagcg gtcgtgtagt tatttttatg tcataagtgg ataatttgtt 60
    agctcctata acaaaagtct gttgcttgtg tttcacattt tggatttcct aatataatgt 120
    tctcttttt   129

    <210> 28
    <211> 128
    <212> DNA
    <213> Mus musculus

    <400> 28
    aattaaaatg tgtcttagtt attgtgccat tatttttatg tcatcactgg ataatatatt 60
    agtgcttagt atcagaaata gtccttatgc tttgtgtttt gaagttccta atgcaatgtt 120
    ctctttct   128

    <210> 29
    <211> 234
    <212> DNA
    <213> Homo sapiens

    <400> 29
    tattctgcct gcttaaatat tactttcctc aaaaagagaa aacaaaaatg ctagattta 60
    ctttatgact tgaatgatgt ggtaatgtcg aactctagta tttagaatta gaatgtttct 120
    tagcggtcgt gtagttattt ttatgtcata agtggataat ttgttagctc ctataacaaa 180
    agtctgttgc ttgtgtttca cattttggat ttcctaatat aatgttctct tttt   234

    <210> 30
    <211> 233
    <212> DNA
    <213> Mus musculus

    <400> 30
    atgtgtttgg cattatgtga agcccaaaca ctaaaaaagg agaacaaaca aaagcgcaga 60
    ctttaaaact caagtggttt ggtaatgtac gactctactg tttagaatta aaatgtgtct 120
    tagttattgt gccattattt ttatgtcatc actggataat atattagtgc ttagtatcag 180
    aaatagtcct tatgctttgt gttttgaagt tcctaatgca atgttctctt tct   233